

What is claimed is:

1. A method for SNP typing which comprises the steps of:
simultaneously amplifying a plurality of nucleotide sequences comprising at least one or more sites of single nucleotide polymorphism using genomic DNA and a plurality of primer pairs; and
typing for distinguishing the site(s) of single nucleotide polymorphism of nucleotides contained in a plurality of nucleotide sequences amplified in the above amplification step using the amplified nucleotide sequences.
2. The method for SNP typing according to claim 1, wherein said step of amplifying employs the polymerase chain reaction using a hot start method.
3. The method for SNP typing according to claim 1, wherein said step of amplifying employs 50 pairs or more primers.
4. The method for SNP typing according to claim 1, wherein said step of typing employs an Invader assay or a TaqMan PCR method.
5. The method for SNP typing according to claim 1, wherein amplification is carried out from 10 ng to 40 ng of genomic DNA.